
Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: markspencer

Timestamp: [year=2009; month=3; day=12; hr=11; min=2; sec=14; ms=883;]

Reviewer Comments:

E287 Invalid WIPO ST.2 date format; Use (YYYY-MM-DD) in <141> E287 Invalid WIPO ST.2 date format; Use (YYYY-MM-DD) in <151> E287 Invalid WIPO ST.2 date format; Use (YYYY-MM-DD) in <151>

<140> 10/565,646

<141> March 24, 2006

<150> PCT/FR2004/001952

<151> July 22, 2004

<150> FR/09161

<151> July 25, 2003

Please change the format of the dates above to (YYYY-MM-DD).

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<211> 648

<212> PRT

<213> Bos taurus

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Ser Gln Ser Tyr Val Pro Leu Ala His Ser Ser Ser Ala Phe Thr Ile

195 200 205

Thr Asp Gln Val Pro Phe Ser Val Ser Val Ser Gln Leu Gln Ala Leu 210 220

Asp Gly Arg Asn Lys Arg Phe Leu Arg Lys Gln Pro Leu Thr Phe Ala 225 230 235 240

"The enumeration of amino acids shall start at the first amino acid as number 1. It shall be marked below the sequence every 5 amino acids. SEQ ID # 6 amino acid number 210 is not in the correct location. Please make all necessary changes.

Validated By CRFValidator v 1.0.3

Application No: 10565646 Version No: 2.0

Input Set:

Output Set:

Started: 2009-02-05 20:45:13.750

Finished: 2009-02-05 20:47:46.221

Elapsed: 0 hr(s) 2 min(s) 32 sec(s) 471 ms

Total Warnings: 21

Total Errors: 335

No. of SeqIDs Defined: 11

Error code		Error Description
E	287	Invalid WIPO ST.2 date format; Use (YYYY-MM-DD) in <141>
E	287	Invalid WIPO ST.2 date format; Use (YYYY-MM-DD)in <151>
E	287	Invalid WIPO ST.2 date format; Use (YYYY-MM-DD)in <151>
E	201	Mandatory field data missing in <223> in SEQ ID (1)
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Ε	320	Wrong Nucleic Acid Designator, ga in SEQID (1)
E	320	Wrong Nucleic Acid Designator, gt in SEQID (1)
E	320	Wrong Nucleic Acid Designator, gg in SEQID (1)
E	320	Wrong Nucleic Acid Designator, ac in SEQID (1)
E	320	Wrong Nucleic Acid Designator, gc in SEQID (1)
Ε	323	Invalid/missing amino acid numbering SEQID (1) POS (6764)

Output Set:

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Elapsed: 0 hr(s) 2 min(s) 32 sec(s) 471 ms

Total Warnings: 21
Total Errors: 335
No. of SeqIDs Defined: 11

Error o	code	Error Description
E 32	20	Wrong Nucleic Acid Designator, gg in SEQID (1)
E 32	20	Wrong Nucleic Acid Designator, gg in SEQID (1)
E 32	20	Wrong Nucleic Acid Designator, ag in SEQID (1)
E 20	01	Mandatory field data missing in <223> in SEQ ID (3)
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E 20	01	Mandatory field data missing in <223> in SEQ ID (3)
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E 20	01	Mandatory field data missing in $<223>$ in SEQ ID (3) This error has occured more than 20 times, will not be displayed
E 32	20	Wrong Nucleic Acid Designator, ga in SEQID (3)
E 32	20	Wrong Nucleic Acid Designator, gt in SEQID (3)
E 32	20	Wrong Nucleic Acid Designator, gg in SEQID (3)
E 32	22	CDS location out of range SEQID (3) At Protien count (157)
E 25	59	Found undefined lettercode; POS (4557) SEQID(3)
E 25	59	Found undefined lettercode; POS (4558) SEQID(3)
E 25	59	Found undefined lettercode; POS (4559) SEQID(3)
E 25	59	Found undefined lettercode; POS (4560) SEQID(3)
E 25	54	The total number of bases conflicts with running total, Input: 4616, Calculated: 4620 SEQID(3)

Output Set:

Started: 2009-02-05 20:45:13.750

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Total Warnings: 21

Total Errors: 335

No. of SeqIDs Defined: 11

Actual SeqID Count: 11

Err	or code	Error Description	
E	254	The total number of bases conflicts with running total, Input: 4676, Calculated: 4680 SEQID(3)	
E	254	The total number of bases conflicts with running total, Input: 4734, Calculated: 4738 SEQID(3)	
E	254	The total number of bases conflicts with running total, Input: 4782, Calculated: 4786 SEQID(3)	:
E	254	The total number of bases conflicts with running total, Input: 4830, Calculated: 4834 SEQID(3)	
E	254	The total number of bases conflicts with running total, Input: 4878, Calculated: 4882 SEQID(3)	
E	254	The total number of bases conflicts with running total, Input: 4926, Calculated: 4930 SEQID(3)	:
Ε	254	The total number of bases conflicts with running total, Input: 4974, Calculated: 4978 SEQID(3)	
Е	254	The total number of bases conflicts with running total, Input: 5022, Calculated: 5026 SEQID(3)	
E	254	The total number of bases conflicts with running total, Input: 5070, Calculated: 5074 SEQID(3)	
E	254	The total number of bases conflicts with running total, Input: 5118, Calculated: 5122 SEQID(3)	
E	254	The total number of bases conflicts with running total, Input: 5166, Calculated: 5170 SEQID(3)	:
Ε	254	The total number of bases conflicts with running total, Input: 5214, Calculated: 5218 SEQID(3)	
Ε	254	The total number of bases conflicts with running total, Input: 5262, Calculated: 5266 SEQID(3)	
E	254	The total number of bases conflicts with running total, Input: 5310, Calculated: 5314 SEQID(3)	:
Е	254	The total number of bases conflicts with running total, Input: 5358, Calculated: 5362 SEQID(3)	

Output Set:

Started: 2009-02-05 20:45:13.750

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Elapsed: 0 hr(s) 2 min(s) 32 sec(s) 471 ms

Total Warnings: 21

Total Errors: 335

No. of SeqIDs Defined: 11

Err	or code	Error Description
E	254	The total number of bases conflicts with running total, Input: 5406, Calculated: 5410 SEQID(3)
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E	254	The total number of bases conflicts with running total, Input: 5523, Calculated: 5527 SEQID(3)
E	254	The total number of bases conflicts with running total, Input: 5583, Calculated: 5587 SEQID(3) This error has occured more than 20 times, will not be displayed
E	320	Wrong Nucleic Acid Designator, gc in SEQID (3)
E	322	CDS location out of range SEQID (3) At Protien count (476)
E	323	Invalid/missing amino acid numbering SEQID (3) POS (6768)
E	322	CDS location out of range SEQID (3) At Protien count (505)
E	320	Wrong Nucleic Acid Designator, gg in SEQID (3)
E	322	CDS location out of range SEQID (3) At Protien count (573)
E	320	Wrong Nucleic Acid Designator, gg in SEQID (3)
E	322	CDS location out of range SEQID (3) At Protien count (603)
E	320	Wrong Nucleic Acid Designator, ag in SEQID (3)
E	322	CDS location out of range SEQID (3) At Protien count (649)
E	253	The number of bases differs from <211> Input: 8146
E	323	Invalid/missing amino acid numbering SEQID (5) POS (2323)
E	320	Wrong Nucleic Acid Designator, ga in SEQID (5)
E	320	Wrong Nucleic Acid Designator, gt in SEQID (5)
E	323	Invalid/missing amino acid numbering SEQID (5) POS (4694)
E	320	Wrong Nucleic Acid Designator, ac in SEQID (5)

Output Set:

Started: 2009-02-05 20:45:13.750 **Finished:** 2009-02-05 20:47:46.221

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Total Warnings: 21
Total Errors: 335
No. of SeqIDs Defined: 11

Error code	Error Description
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E 320	Wrong Nucleic Acid Designator, ag in SEQID (5) This error has occured more than 20 times, will not be displayed
W 112	Upper case found in data; Found at position(0) SeqId(6)
W 112	Upper case found in data; Found at position(3) SeqId(6)
E 259	Found undefined lettercode; POS (6) SEQID(6)
W 112	Upper case found in data; Found at position(6) SeqId(6)
E 259	Found undefined lettercode; POS (8) SEQID(6)
E 342	'n' position not defined found at POS: 9 SEQID(6)
W 112	Upper case found in data; Found at position(9) SeqId(6)
E 259	Found undefined lettercode; POS (12) SEQID(6)
W 112	Upper case found in data; Found at position(12) SeqId(6)
E 259	Found undefined lettercode; POS (15) SEQID(6)
W 112	Upper case found in data; Found at position(15) SeqId(6)
E 259	Found undefined lettercode; POS (18) SEQID(6)
W 112	Upper case found in data; Found at position(18) SeqId(6)
E 259	Found undefined lettercode; POS (20) SEQID(6)
W 112	Upper case found in data; Found at position(21) SeqId(6)
E 259	Found undefined lettercode; POS (24) SEQID(6)
W 112	Upper case found in data; Found at position(24) SeqId(6)
E 259	Found undefined lettercode; POS (26) SEQID(6)
W 112	Upper case found in data; Found at position(27) SeqId(6)
E 259	Found undefined lettercode; POS (30) SEQID(6)

Output Set:

Started: 2009-02-05 20:45:13.750 **Finished:** 2009-02-05 20:47:46.221

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Total Warnings: 21
Total Errors: 335
No. of SeqIDs Defined: 11

Err	ror code	Error Description
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E	259	Found undefined lettercode; POS (32) SEQID(6)
W	112	Upper case found in data; Found at position(33) SeqId(6)
E	259	Found undefined lettercode; POS (35) SEQID(6)
E	342	'n' position not defined found at POS: 36 SEQID(6)
W	112	Upper case found in data; Found at position(36) SeqId(6)
E	259	Found undefined lettercode; POS (38) SEQID(6)
W	112	Upper case found in data; Found at position(39) SeqId(6)
E	259	Found undefined lettercode; POS (41) SEQID(6)
E	342	'n' position not defined found at POS: 42 SEQID(6)
W	112	Upper case found in data; Found at position(42) SeqId(6)
E	259	Found undefined lettercode; POS (44) SEQID(6)
W	112	Upper case found in data; Found at position(45) SeqId(6)
E	259	Found undefined lettercode; POS (47) SEQID(6)
E	259	Found undefined lettercode; POS (49) SEQID(6) This error has occured more than 20 times, will not be displayed
E	323	Invalid/missing amino acid numbering SEQID (6) POS (209)
E	323	Invalid/missing amino acid numbering SEQID (6)at Protein (210)
E	323	Invalid/missing amino acid numbering SEQID (6) POS (214)
E	323	Invalid/missing amino acid numbering SEQID (6)at Protein (215)
E	323	Invalid/missing amino acid numbering SEQID (6) POS (219)
E	323	Invalid/missing amino acid numbering SEQID (6)at Protein (220)
E	323	Invalid/missing amino acid numbering SEQID (6) POS (224)

Output Set:

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Total Warnings: 21
Total Errors: 335
No. of SeqIDs Defined: 11

Error code		Error Description
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E	323	Invalid/missing amino acid numbering SEQID (6) POS (229)
E	323	Invalid/missing amino acid numbering SEQID (6)at Protein (230)
E	323	Invalid/missing amino acid numbering SEQID (6) POS (234)
Ε	323	Invalid/missing amino acid numbering SEQID (6)at Protein (235)
Ε	323	Invalid/missing amino acid numbering SEQID (6) POS (239)
Ε	323	Invalid/missing amino acid numbering SEQID (6)at Protein (240)
Ε	323	Invalid/missing amino acid numbering SEQID (6) POS (244)
Ε	323	Invalid/missing amino acid numbering SEQID (6) at Protein (245) This error has occured more than 20 times, will not be displayed
E	331	Count of Protein differs from the <211> tag Input: 648
W	213	Artificial or Unknown found in <213> in SEQ ID (7)
W	213	Artificial or Unknown found in <213> in SEQ ID (8)
W	213	Artificial or Unknown found in <213> in SEQ ID (9)
W	213	Artificial or Unknown found in <213> in SEQ ID (10)
W	213	Artificial or Unknown found in <213> in SEQ ID (11)

SEQUENCE LISTING

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Lafort, Marie-Piere,
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Gly Leu Ser Ile Gly Thr Asp Lys Ala Met Leu Gly Thr Tyr Asn Met	
Gly Leu Ser Ile Gly Thr Asp Lys Ala Met Leu Gly Thr Tyr Asn Met 170 180	4445
Gly Leu Ser Ile Gly Thr Asp Lys Ala Met Leu Gly Thr Tyr Asn Met 170 175 180 gaa gtg act gtc tac cac cgc cgg ggg tcc cag agc tat gtg ccc ctc	4445
Gly Leu Ser Ile Gly Thr Asp Lys Ala Met Leu Gly Thr Tyr Asn Met 170 175 180 gaa gtg act gtc tac cac cgc cgg ggg tcc cag agc tat gtg ccc ctc Glu Val Thr Val Tyr His Arg Arg Gly Ser Gln Ser Tyr Val Pro Leu	4445
Gly Leu Ser Ile Gly Thr Asp Lys Ala Met Leu Gly Thr Tyr Asn Met 170 175 180 gaa gtg act gtc tac cac cgc cgg ggg tcc cag agc tat gtg ccc ctc	4445
Gly Leu Ser Ile Gly Thr Asp Lys Ala Met Leu Gly Thr Tyr Asn Met 170	
Gly Leu Ser Ile Gly Thr Asp Lys Ala Met Leu Gly Thr Tyr Asn Met 170 175 180 gaa gtg act gtc tac cac cgc cgg ggg tcc cag agc tat gtg ccc ctc Glu Val Thr Val Tyr His Arg Arg Gly Ser Gln Ser Tyr Val Pro Leu 185 190 195 200 gct cac tcc agt tca gcc ttc acc att act g gtaaggactg aggagggac	4445
Gly Leu Ser Ile Gly Thr Asp Lys Ala Met Leu Gly Thr Tyr Asn Met 170	
Gly Leu Ser Ile Gly Thr Asp Lys Ala Met Leu Gly Thr Tyr Asn Met 170 175 180 gaa gtg act gtc tac cac cgc cgg ggg tcc cag agc tat gtg ccc ctc Glu Val Thr Val Tyr His Arg Arg Gly Ser Gln Ser Tyr Val Pro Leu 185 190 195 200 gct cac tcc agt tca gcc ttc acc att act g gtaaggactg aggagggac	
Gly Leu Ser Ile Gly Thr Asp Lys Ala Met Leu Gly Thr Tyr Asn Met 170	4496
Gly Leu Ser Ile Gly Thr Asp Lys Ala Met Leu Gly Thr Tyr Asn Met 170 175 180 gaa gtg act gtc tac cac cgc cgg ggg tcc cag agc tat gtg ccc ctc Glu Val Thr Val Tyr His Arg Arg Gly Ser Gln Ser Tyr Val Pro Leu 185 190 195 200 gct cac tcc agt tca gcc ttc acc att act g gtaaggactg aggaggggac Ala His Ser Ser Ser Ala Phe Thr Ile Thr 205 210 aaggccagtt gcagggcagg agaaggtggg gaggctgggc tggacaggaa aggggaaaga	4496
Gly Leu Ser Ile Gly Thr Asp Lys Ala Met Leu Gly Thr Tyr Asn Met 170 175 180 gaa gtg act gtc tac cac cgc cgg ggg tcc cag agc tat gtg ccc ctc Glu Val Thr Val Tyr His Arg Arg Gly Ser Gln Ser Tyr Val Pro Leu 185 190 195 200 gct cac tcc agt tca gcc ttc acc att act g gtaaggactg aggaggggac Ala His Ser Ser Ser Ala Phe Thr Ile Thr 205 210 aaggccagtt gcagggcagg agaaggtggg gaggctgggc tggacaggaa aggggaaaga ggaaatggtg tgtaacctta caggggcaga accaggaaga tgtgggcaga gggatgtggg	4496 4556 4616
Gly Leu Ser Ile Gly Thr Asp Lys Ala Met Leu Gly Thr Tyr Asn Met 170 175 180 gaa gtg act gtc tac cac cgc cgg ggg tcc cag agc tat gtg ccc ctc Glu Val Thr Val Tyr His Arg Arg Gly Ser Gln Ser Tyr Val Pro Leu 185 190 195 200 gct cac tcc agt tca gcc ttc acc att act g gtaaggactg aggaggggac Ala His Ser Ser Ser Ala Phe Thr Ile Thr 205 210 aaggccagtt gcagggcagg agaaggtggg gaggctgggc tggacaggaa aggggaaaga	4496

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cag gtg ccc ttc tct gtg agt gtg tct cag ctg cag gcc ttg gat gga Gln Val Pro Phe Ser Val Ser Val Ser Gln Leu Gln Ala Leu Asp Gly 215 220 225	
agg aac aag cgc ttc ctg aga aag cag cct ctg acc ttt gcc ctc cag Arg Asn Lys Arg Phe Leu Arg Lys Gln Pro Leu Thr Phe Ala Leu Gln 230 235 240	
ctc cat gat ccc agt ggc tat ttg gct ggg gct gac ctt tcc tac acc Leu His Asp Pro Ser Gly Tyr Leu Ala Gly Ala Asp Leu Ser Tyr Thr 245 250 255	
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cct gag cct gcg ggt tca aat act agc tca ttc atg cct aca gaa ggt Pro Glu Pro Ala Gly Ser Asn Thr Ser Ser Phe Met Pro Thr Glu Gly 420 425 430 430	5406
act gca g gtaagggggc caccatgaat gagttcatag aggtggggca tttgtcacag Thr Ala	5463
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Leu Pro Lys Glu Ala 510

tgc atg gac atc tca tcg cca ggg tgt cag ctg cct gcc cag cgg ctg Cys Met Asp Ile Ser Ser Pro Gly Cys Gln Leu Pro Ala Gln Arg Leu 515 520 525	6938
tgt cag cct gtg ccc ccc agc cca gcc tgc cag ctg gtt ttg cac cag Cys Gln Pro Val Pro Pro Ser Pro Ala Cys Gln Leu Val Leu His Gln 530 535 540	6986
gta ctg aag ggt ggc tca ggg acc tac tgc ctc aat gtg tct ttg gct Val Leu Lys Gly Gly Ser Gly Thr Tyr Cys Leu Asn Val Ser Leu Ala 545 550 555	7034
gat gcc aat agc ctg gcg atg gtc agc acc cag ctt gtc atg cct g Asp Ala Asn Ser Leu Ala Met Val Ser Thr Gln Leu Val Met Pro 560 565 570	7080
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